

*Brancock. M*

# ENTERED



1600

## RAW SEQUENCE LISTING

DATE: 04/22/2003

PATENT APPLICATION: US/09/848,664A

TIME: 14:56:52

Input Set : N:\Crf3\RULE60\09848664.raw.txt

Output Set: N:\CRF4\04222003\I848664A.raw

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1 <110> APPLICANT: Sakiyama-Elbert, Shelly E.
2   Hubbell, Jeffrey A.
3 <120> TITLE OF INVENTION: Controlled Release of Non-Heparin Binding Growth
4   Factors from Heparin Containing Matrices
5 <130> FILE REFERENCE: ETH 108
6 <140> CURRENT APPLICATION NUMBER: US/09/848,664A
7 <141> CURRENT FILING DATE: 2001-05-03
8 <150> PRIOR APPLICATION NUMBER: US/09/298,084A
9 <151> PRIOR FILING DATE: 1999-04-22
10 <160> NUMBER OF SEQ ID NOS: 31
11 <170> SOFTWARE: PatentIn Ver. 2.1
13 <210> SEQ ID NO: 1
14 <211> LENGTH: 14
15 <212> TYPE: PRT
16 <213> ORGANISM: Homo sapiens
17 <220> FEATURE:
18 <221> NAME/KEY: MOD_RES
19 <222> LOCATION: (2)
20 <223> OTHER INFORMATION: Xaa is bAla (Beta Alanine)
21 <400> SEQUENCE: 1
W--> 22   Lys Xaa Phe Ala Lys Leu Ala Ala Arg Leu Tyr Arg Lys Ala
23         1             5             10
25 <210> SEQ ID NO: 2
26 <211> LENGTH: 8
27 <212> TYPE: PRT
28 <213> ORGANISM: Homo sapiens
29 <400> SEQUENCE: 2
30   Tyr Lys Lys Ile Ile Lys Lys Leu
31         1             5
33 <210> SEQ ID NO: 3
34 <211> LENGTH: 14
35 <212> TYPE: PRT
36 <213> ORGANISM: Homo sapiens
37 <400> SEQUENCE: 3
38   Lys His Lys Gly Arg Asp Val Ile Leu Lys Lys Asp Val Arg
39         1             5             10
41 <210> SEQ ID NO: 4
42 <211> LENGTH: 14
43 <212> TYPE: PRT
44 <213> ORGANISM: Homo sapiens
45 <220> FEATURE:
46 <221> NAME/KEY: MOD_RES
47 <222> LOCATION: (2)

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48 <223> OTHER INFORMATION: Xaa is bALA (Beta Alanine)
49 <400> SEQUENCE: 4
W--> 50      Arg Xaa Phe Ala Arg Leu Ala Ala Arg Leu Tyr Arg Arg Ala
51          1              5              10
53 <210> SEQ ID NO: 5
54 <211> LENGTH: 12
55 <212> TYPE: PRT
56 <213> ORGANISM: Homo sapiens
57 <400> SEQUENCE: 5
58      Lys Asp Pro Lys Arg Leu Tyr Arg Ser Arg Lys Tyr
59          1              5              10
61 <210> SEQ ID NO: 6
62 <211> LENGTH: 11
63 <212> TYPE: PRT
64 <213> ORGANISM: Homo sapiens
65 <400> SEQUENCE: 6
66      Cys Val Leu Ser Arg Lys Ala Val Arg Arg Ala
67          1              5              10
69 <210> SEQ ID NO: 7
70 <211> LENGTH: 10
71 <212> TYPE: PRT
72 <213> ORGANISM: Homo sapiens
73 <400> SEQUENCE: 7
74      Cys Ala Leu Ser Arg Lys Ile Gly Arg Thr
75          1              5              10
77 <210> SEQ ID NO: 8
78 <211> LENGTH: 9
79 <212> TYPE: PRT
80 <213> ORGANISM: Homo sapiens
81 <400> SEQUENCE: 8
82      Cys Thr Leu Thr Ile Lys Arg Gly Arg
83          1              5
85 <210> SEQ ID NO: 9
86 <211> LENGTH: 70
87 <212> TYPE: PRT
88 <213> ORGANISM: Homo sapiens
89 <400> SEQUENCE: 9
90      Ala Leu Asp Thr Asn Tyr Cys Phe Ser Ser Thr Glu Lys Asn Cys Cys
91          1              5              10              15
92      Val Arg Gln Leu Tyr Ile Asp Phe Arg Lys Asp Leu Gly Trp Lys Trp
93          20              25              30
94      Ile His Glu Pro Lys Gly Tyr His Ala Asn Phe Cys Leu Gly Pro Cys
95          35              40              45
96      Pro Tyr Ile Trp Ser Leu Asp Thr Gln Tyr Ser Lys Val Leu Ala Leu
97          50              55              60
98      Tyr Asn Gln His Asn Pro
99          65              70
101 <210> SEQ ID NO: 10
102 <211> LENGTH: 70

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103 <212> TYPE: PRT
104 <213> ORGANISM: Homo sapiens
105 <400> SEQUENCE: 10
106   Ala Leu Asp Ala Ala Tyr Cys Phe Arg Asn Val Gln Asp Asn Cys Cys
107       1             5             10             15
108   Leu Arg Pro Leu Tyr Ile Asp Phe Lys Arg Asp Leu Gly Trp Lys Trp
109               20             25             30
110   Ile His Glu Pro Lys Gly Tyr Asn Ala Asn Phe Cys Ala Gly Ala Cys
111               35             40             45
112   Pro Tyr Leu Trp Ser Ser Asp Thr Gln His Ser Arg Val Leu Ser Leu
113       50             55             60
114   Tyr Asn Thr Ile Asn Pro
115       65             70
117 <210> SEQ ID NO: 11
118 <211> LENGTH: 70
119 <212> TYPE: PRT
120 <213> ORGANISM: Homo sapiens
121 <400> SEQUENCE: 11
122   Ala Leu Asp Thr Asn Tyr Cys Phe Arg Asn Leu Glu Glu Asn Cys Cys
123       1             5             10             15
124   Val Arg Pro Leu Tyr Ile Asp Phe Arg Gln Asp Leu Gly Trp Lys Trp
125               20             25             30
126   Val His Glu Pro Lys Gly Tyr Tyr Ala Asn Phe Cys Ser Gly Pro Cys
127               35             40             45
128   Pro Tyr Leu Arg Ser Ala Asp Thr Thr His Ser Thr Val Leu Gly Leu
129       50             55             60
130   Tyr Asn Thr Leu Asn Pro
131       65             70
133 <210> SEQ ID NO: 12
134 <211> LENGTH: 42
135 <212> TYPE: PRT
136 <213> ORGANISM: Homo sapiens
137 <400> SEQUENCE: 12
138   Gly Ala Ser Ala Ala Pro Cys Cys Val Pro Gln Ala Leu Glu Pro Leu
139       1             5             10             15
140   Pro Ile Val Tyr Tyr Val Gly Arg Lys Pro Lys Val Glu Gln Leu Ser
141               20             25             30
142   Asn Met Ile Val Arg Ser Cys Lys Cys Ser
143       35             40
145 <210> SEQ ID NO: 13
146 <211> LENGTH: 42
147 <212> TYPE: PRT
148 <213> ORGANISM: Homo sapiens
149 <400> SEQUENCE: 13
150   Glu Ala Ser Ala Ser Pro Cys Cys Val Ser Gln Asp Leu Glu Pro Leu
151       1             5             10             15
152   Thr Ile Leu Tyr Tyr Ile Gly Lys Thr Pro Lys Ile Glu Gln Leu Ser
153               20             25             30
154   Asn Met Ile Val Lys Ser Cys Lys Cys Ser

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Input Set : N:\Crf3\RULE60\09848664.raw.txt

Output Set: N:\CRF4\04222003\I848664A.raw

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155          35          40
157 <210> SEQ ID NO: 14
158 <211> LENGTH: 42
159 <212> TYPE: PRT
160 <213> ORGANISM: Homo sapiens
161 <400> SEQUENCE: 14
162      Glu Ala Ser Ala Ser Pro Cys Cys Val Pro Gln Asp Leu Glu Pro Leu
163          1          5          10          15
164      Thr Ile Leu Tyr Tyr Val Gly Arg Thr Pro Lys Val Glu Gln Leu Ser
165          20          25          30
166      Asn Met Val Val Lys Ser Cys Lys Cys Ser
167          35          40
169 <210> SEQ ID NO: 15
170 <211> LENGTH: 294
171 <212> TYPE: PRT
172 <213> ORGANISM: Homo sapiens
173 <400> SEQUENCE: 15
174      Phe Ser Gln Ser Phe Arg Glu Val Ala Gly Arg Phe Leu Ala Ser Glu
175          1          5          10          15
176      Ala Ser Thr His Leu Leu Val Phe Gly Met Glu Gln Arg Leu Pro Pro
177          20          25          30
178      Asn Ser Glu Leu Val Gln Ala Val Leu Arg Leu Phe Gln Glu Pro Val
179          35          40          45
180      Pro Gln Gly Ala Leu His Arg His Gly Arg Leu Ser Pro Ala Ala Pro
181          50          55          60
182      Lys Ala Arg Val Thr Val Glu Trp Leu Val Arg Asp Asp Gly Ser Asn
183          65          70          75          80
184      Arg Thr Ser Leu Ile Asp Ser Arg Leu Val Ser Val His Glu Ser Gly
185          85          90          95
186      Trp Lys Ala Phe Asp Val Thr Glu Ala Val Asn Phe Trp Gln Gln Leu
187          100          105          110
188      Ser Arg Pro Pro Glu Pro Leu Leu Val Gln Val Ser Val Gln Arg Glu
189          115          120          125
190      His Leu Gly Pro Leu Ala Ser Gly Ala His Lys Leu Val Arg Phe Ala
191          130          135          140
192      Ser Gln Gly Ala Pro Ala Gly Leu Gly Glu Pro Gln Leu Glu Leu His
193          145          150          155          160
194      Thr Leu Asp Leu Arg Asp Tyr Gly Ala Gln Gly Asp Cys Asp Pro Glu
195          165          170          175
196      Ala Pro Met Thr Glu Gly Thr Arg Cys Cys Arg Gln Glu Met Tyr Ile
197          180          185          190
198      Asp Leu Gln Gly Met Lys Trp Ala Lys Asn Trp Val Leu Glu Pro Pro
199          195          200          205
200      Gly Phe Leu Ala Tyr Glu Cys Val Gly Thr Cys Gln Gln Pro Pro Glu
201          210          215          220
202      Ala Leu Ala Phe Asn Trp Pro Phe Leu Gly Pro Arg Gln Cys Ile Ala
203          225          230          235          240
204      Ser Glu Thr Ala Ser Leu Pro Met Ile Val Ser Ile Lys Glu Gly Gly
205          245          250          255

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206      Arg Thr Arg Pro Gln Val Val Ser Leu Pro Asn Met Arg Val Gln Lys
207                260                265                270
208      Cys Ser Cys Ala Ser Asp Gly Ala Leu Val Pro Arg Arg Leu Gln His
209                275                280                285
210      Arg Pro Trp Cys Ile His
211                290
213 <210> SEQ ID NO: 16
214 <211> LENGTH: 73
215 <212> TYPE: PRT
216 <213> ORGANISM: Homo sapiens
217 <400> SEQUENCE: 16
218      Ser Pro Asp Lys Gln Met Ala Val Leu Pro Arg Arg Glu Arg Asn Arg
219      1                5                10                15
220      Gln Ala Ala Ala Ala Asn Pro Glu Asn Ser Arg Gly Lys Gly Arg Arg
221                20                25                30
222      Gly Gln Arg Gly Lys Asn Arg Gly Cys Val Leu Thr Ala Ile His Leu
223                35                40                45
224      Asn Val Thr Asp Leu Gly Leu Gly Tyr Glu Thr Lys Glu Glu Leu Ile
225                50                55                60
226      Phe Arg Tyr Cys Ser Gly Ser Cys Asp
227      65                70
229 <210> SEQ ID NO: 17
230 <211> LENGTH: 73
231 <212> TYPE: PRT
232 <213> ORGANISM: Homo sapiens
233 <400> SEQUENCE: 17
234      Leu Gly Ala Arg Pro Cys Gly Leu Arg Glu Leu Glu Val Arg Val Ser
235      1                5                10                15
236      Glu Leu Gly Leu Gly Tyr Ala Ser Asp Glu Thr Val Leu Phe Arg Tyr
237                20                25                30
238      Cys Ala Gly Ala Cys Glu Ala Ala Ala Arg Val Tyr Asp Leu Gly Leu
239                35                40                45
240      Arg Arg Leu Arg Gln Arg Arg Arg Leu Arg Arg Glu Arg Val Arg Ala
241                50                55                60
242      Gln Pro Cys Cys Arg Pro Thr Ala Tyr
243      65                70
245 <210> SEQ ID NO: 18
246 <211> LENGTH: 61
247 <212> TYPE: PRT
248 <213> ORGANISM: Homo sapiens
249 <400> SEQUENCE: 18
250      Ala Ala Glu Thr Thr Tyr Asp Lys Ile Leu Lys Asn Leu Ser Arg Asn
251      1                5                10                15
252      Arg Arg Leu Val Ser Asp Lys Val Gly Gln Ala Cys Cys Arg Pro Ile
253                20                25                30
254      Ala Phe Asp Asp Asp Leu Ser Phe Leu Asp Asp Asn Leu Val Tyr His
255                35                40                45
256      Ile Leu Arg Lys His Ser Ala Lys Arg Cys Gly Cys Ile
257      50                55                60

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RAW SEQUENCE LISTING ERROR SUMMARY      DATE: 04/22/2003  
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; Xaa Pos. 2  
Seq#:4; Xaa Pos. 2

**VERIFICATION SUMMARY**

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Input Set : N:\Crf3\RULE60\09848664.raw.txt

Output Set: N:\CRF4\04222003\I848664A.raw

L:22 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0

L:50 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0